

STN Results 09/758,003

(FILE 'HOME' ENTERED AT 15:21:42 ON 19 SEP 2005)

FILE 'BIOSIS, MEDLINE, EMBASE, CAPLUS, SCISEARCH' ENTERED AT  
15:22:07 ON  
19 SEP 2005

L1 12 S BAICHWAL  
L2 0 S HUANG JIANING  
L3 164 S HUANG J  
L4 46 S GOEDDEL

(FILE 'HOME' ENTERED AT 11:22:29 ON 21 SEP 2005)

FILE 'CAPLUS, EMBASE, MEDLINE, SCISEARCH, BIOSIS' ENTERED AT  
11:23:38 ON  
21 SEP 2005

L1 9001 S RIP  
L2 51 S RIP KINASE  
L3 343 S RIP PROTEIN  
L4 4524 S RIP AND PROTEIN  
L5 955 S RIP AND KINASE

09/758,003 Results

SEQ ID NO: 2-oligo search

SUMMARIES

Result	Query					Description
No.	Score	Match	Length	DB	ID	
1	671	100.0	671	3	AAY78502	Aay78502 Human RIP
2	671	100.0	671	4	AAB82091	Aab82091 Human Rec
3	671	100.0	671	4	ABG16302	Abg16302 Novel hum
4	671	100.0	671	5	AAU80370	Aau80370 Human cel
5	671	100.0	671	5	ABB83794	Abb83794 Human RIP
6	671	100.0	671	7	ADP65296	Adp65296 Human rec
7	671	100.0	671	8	ADS88161	Ads88161 Human pro
8	570	84.9	671	2	AAW15461	Aaw15461 Human rec
9	570	84.9	671	7	ADJ70448	Adj70448 Human hea
10	339	50.5	671	2	AAW04628	Aaw04628 Human rec
11	222	33.1	273	7	ADM05095	Adm05095 Human pro
12	153	22.8	948	4	ABG16304	Abg16304 Novel hum
13	98	14.6	99	7	ADC08901	Adc08901 Recombina
14	98	14.6	100	7	ADC08899	Adc08899 Recombina
15	76	11.3	77	6	ADA49713	Ada49713 Death dom

SUMMARIES

Result	Query					Description
No.	Score	Match	Length	DB	ID	
1	671	100.0	671	3	US-09-132-118-2	Sequence 2, Appli
2	671	100.0	671	4	US-09-345-473E-29	Sequence 29, Appl
3	368	54.8	709	1	US-08-444-005-17	Sequence 17, Appl
4	87	13.0	87	3	US-09-042-785A-29	Sequence 29, Appl
5	77	11.5	77	3	US-08-995-159-7	Sequence 7, Appli
6	77	11.5	77	4	US-09-545-605-7	Sequence 7, Appli
7	76	11.3	77	4	US-08-828-683A-27	Sequence 27, Appl
8	42	6.3	42	1	US-08-444-005-25	Sequence 25, Appl
9	39	5.8	39	1	US-08-702-970-4	Sequence 4, Appli
10	38	5.7	38	1	US-08-444-005-19	Sequence 19, Appl
11	26	3.9	42	1	US-08-444-005-24	Sequence 24, Appl
12	26	3.9	656	1	US-08-444-005-15	Sequence 15, Appl
13	26	3.9	656	3	US-09-069-023-28	Sequence 28, Appl
14	26	3.9	656	4	US-09-345-473E-30	Sequence 30, Appl
15	22	3.3	22	1	US-08-444-005-31	Sequence 31, Appl

SUMMARIES

Result	Query					Description
No.	Score	Match	Length	DB	ID	
1	671	100.0	671	9	US-09-758-003-2	Sequence 2, Appli
2	671	100.0	671	9	US-09-862-027-29	Sequence 29, Appl
3	671	100.0	671	10	US-09-981-397A-16	Sequence 16, Appl
4	671	100.0	671	17	US-10-989-228-29	Sequence 29, Appl
5	671	100.0	671	18	US-10-973-858-38	Sequence 38, Appl
6	570	84.9	671	16	US-10-408-765A-2254	Sequence 2254, Ap
7	222	33.1	273	15	US-10-108-260A-3780	Sequence 3780, Ap
8	98	14.6	99	15	US-10-359-439-3	Sequence 3, Appli
9	98	14.6	100	15	US-10-359-439-1	Sequence 1, Appli
10	77	11.5	77	14	US-10-287-594-7	Sequence 7, Appli
11	76	11.3	77	13	US-10-112-793-27	Sequence 27, Appl
12	52	7.7	52	9	US-09-864-761-35037	Sequence 35037, A
13	26	3.9	67	18	US-10-791-513-5	Sequence 5, Appli
14	26	3.9	656	9	US-09-862-027-30	Sequence 30, Appl
15	26	3.9	656	17	US-10-989-228-30	Sequence 30, Appl

No.	Score	Match	Length	DB	ID	Description
1	2016	100.0	2016	3	US-09-132-118-1	Sequence 1, Appli
2	2016	100.0	2617	3	US-09-161-443-1	Sequence 1, Appli
3	1965	97.5	2617	4	US-09-023-655-914	Sequence 914, App
4	1842	91.4	9687	3	US-09-133-944-2	Sequence 2, Appli
5	1842	91.4	9687	3	US-09-208-827-2	Sequence 2, Appli
6	1842	91.4	9687	4	US-10-043-074-2	Sequence 2, Appli
7	1710	84.8	2137	1	US-08-444-005-16	Sequence 16, Appli
8	259	12.8	606	3	US-09-328-111-495	Sequence 495, App
9	136	6.7	366	4	US-09-513-999C-3314	Sequence 3314, Ap
10	133	6.6	529	4	US-09-854-133-676	Sequence 676, App
11	44	2.2	2268	1	US-08-444-005-14	Sequence 14, Appli
12	30	1.5	30	3	US-09-161-443-4	Sequence 4, Appli
13	28	1.4	28	2	US-08-859-998-1135	Sequence 1135, Ap
14	28	1.4	28	3	US-09-225-928-1135	Sequence 1135, Ap
15	28	1.4	28	4	US-09-225-201B-1135	Sequence 1135, Ap

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Result	No.	Query					Description
		Score	Match	Length	DB	ID	
	1	2016	100.0	2016	9	US-09-758-003-1	Sequence 1, Appli
	2	2016	100.0	2617	10	US-09-981-397A-15	Sequence 15, Appli
	3	2016	100.0	2617	22	US-10-973-858-37	Sequence 37, Appli
	4	1965	97.5	2617	18	US-10-641-643-914	Sequence 914, App
	5	1842	91.4	9687	9	US-09-208-827-2	Sequence 2, Appli
	6	1842	91.4	9687	14	US-10-043-074-2	Sequence 2, Appli
	7	671	33.3	2862	17	US-10-108-260A-1337	Sequence 1337, Ap
c	8	311	15.4	331	9	US-09-796-692-6140	Sequence 6140, Ap
c	9	311	15.4	331	14	US-10-040-862-6140	Sequence 6140, Ap
c	10	311	15.4	331	17	US-10-057-475B-6140	Sequence 6140, Ap
c	11	311	15.4	331	17	US-10-154-884B-6140	Sequence 6140, Ap
c	12	311	15.4	331	19	US-10-764-324-6140	Sequence 6140, Ap
	13	262	13.0	392	10	US-09-918-995-17425	Sequence 17425, A
	14	259	12.8	606	9	US-09-879-536-495	Sequence 495, App
	15	157	7.8	193	9	US-09-864-761-18496	Sequence 18496, A

SEQ ID NO: 1

#### SUMMARIES

Result	No.	Query					Description
		Score	Match	Length	DB	ID	
	1	2016	100.0	2016	6	AR145196	AR145196 Sequence
	2	2016	100.0	2617	6	AX429238	AX429238 Sequence
	3	2016	100.0	2617	6	AX460995	AX460995 Sequence
	4	2016	100.0	2617	9	HSU50062	U50062 Homo sapien
	5	2014.4	99.9	2016	6	CQ720549	CQ720549 Sequence
	6	2014.4	99.9	2617	6	AR380369	AR380369 Sequence
	7	2006.4	99.5	2137	6	I68123	I68123 Sequence 16
	8	2000.4	99.2	9687	6	AR165919	AR165919 Sequence
	9	2000.4	99.2	9687	6	AR429967	AR429967 Sequence
	10	1215.8	60.3	2268	6	I68122	I68122 Sequence 14
	11	1215.8	60.3	2268	10	MMU25995	U25995 Mus musculu
	12	1215.8	60.3	3225	10	BC058162	BC058162 Mus muscu
	13	1215.8	60.3	4512	10	BC050905	BC050905 Mus muscu
	14	1215.8	60.3	4512	10	BC054542	BC054542 Mus muscu
	15	1114.2	55.3	1240	9	HSU25994	U25994 Human cell

#### SUMMARIES

Result	No.	Score	Match	Length	DB	ID	Description

1	2016	100.0	2016	4	AAF86480	Aaf86480 Human Rec
2	2016	100.0	2016	5	AAS80489	Aas80489 DNA encod
3	2016	100.0	2016	11	ADP65098	Adp65098 Human rec
4	2016	100.0	2617	3	AAZ89748	Aaz89748 Human RIP
5	2016	100.0	2617	6	ABK51170	Abk51170 cDNA enco
6	2016	100.0	2617	6	ABN81397	Abn81397 Human RIP
7	2016	100.0	2617	11	ADP65671	Adp65671 Human RIP
8	2014.4	99.9	2016	2	AAT66408	Aat66408 Human rec
9	2014.4	99.9	2617	11	ADI31588	Adi31588 Human cDN
10	2003.2	99.4	2137	2	AAT43753	Aat43753 Human rec
11	2000.4	99.2	9687	4	AAS13810	Aas13810 DNA encod
12	2000.4	99.2	9687	6	ABK15129	Abk15129 Plasmid p
13	2000.4	99.2	9687	8	ABX95502	Abx95502 Novel shu
14	1215.8	60.3	2268	2	AAT43752	Aat43752 Mouse rec
15	1215.8	60.3	2268	2	AAV99827	Aav99827 Mouse rec

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	No.	Score	Match	Length	DB	ID
1	2016	100.0	2016	3	US-09-132-118-1	Sequence 1, Appli
2	2016	100.0	2617	3	US-09-161-443-1	Sequence 1, Appli
3	2014.4	99.9	2617	4	US-09-023-655-914	Sequence 914, App
4	2006.4	99.5	2137	1	US-08-444-005-16	Sequence 16, Appli
5	2000.4	99.2	9687	3	US-09-133-944-2	Sequence 2, Appli
6	2000.4	99.2	9687	3	US-09-208-827-2	Sequence 2, Appli
7	2000.4	99.2	9687	4	US-10-043-074-2	Sequence 2, Appli
8	1215.8	60.3	2268	1	US-08-444-005-14	Sequence 14, Appli
9	445	22.1	529	4	US-09-854-133-676	Sequence 676, App
10	302.4	15.0	606	3	US-09-328-111-495	Sequence 495, App
11	217.2	10.8	366	4	US-09-513-999C-3314	Sequence 3314, Ap
12	71.2	3.5	3699	4	US-09-949-016-759	Sequence 759, App
13	71.2	3.5	3806	4	US-09-949-016-1044	Sequence 1044, Ap
14	69.6	3.5	3542	4	US-09-949-016-5013	Sequence 5013, Ap
15	69.6	3.5	3542	4	US-09-949-016-5014	Sequence 5014, Ap

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1	2016	100.0	2016	9	US-09-758-003-1	Sequence 1, Appli
2	2016	100.0	2617	10	US-09-981-397A-15	Sequence 15, Appli
3	2016	100.0	2617	22	US-10-973-858-37	Sequence 37, Appli
4	2014.4	99.9	2617	18	US-10-641-643-914	Sequence 914, App
5	2000.4	99.2	9687	9	US-09-208-827-2	Sequence 2, Appli
6	2000.4	99.2	9687	14	US-10-043-074-2	Sequence 2, Appli
7	920	45.6	2862	17	US-10-108-260A-1337	Sequence 1337, Ap
8	445	22.1	529	9	US-09-854-133-676	Sequence 676, App
9	445	22.1	529	15	US-10-144-649A-676	Sequence 676, App
10	382.4	19.0	392	10	US-09-918-995-17425	Sequence 17425, A
c 11	311	15.4	331	9	US-09-796-692-6140	Sequence 6140, Ap
c 12	311	15.4	331	14	US-10-040-862-6140	Sequence 6140, Ap
c 13	311	15.4	331	17	US-10-057-475B-6140	Sequence 6140, Ap
c 14	311	15.4	331	17	US-10-154-884B-6140	Sequence 6140, Ap
c 15	311	15.4	331	19	US-10-764-324-6140	Sequence 6140, Ap